

Fig. 1a-f

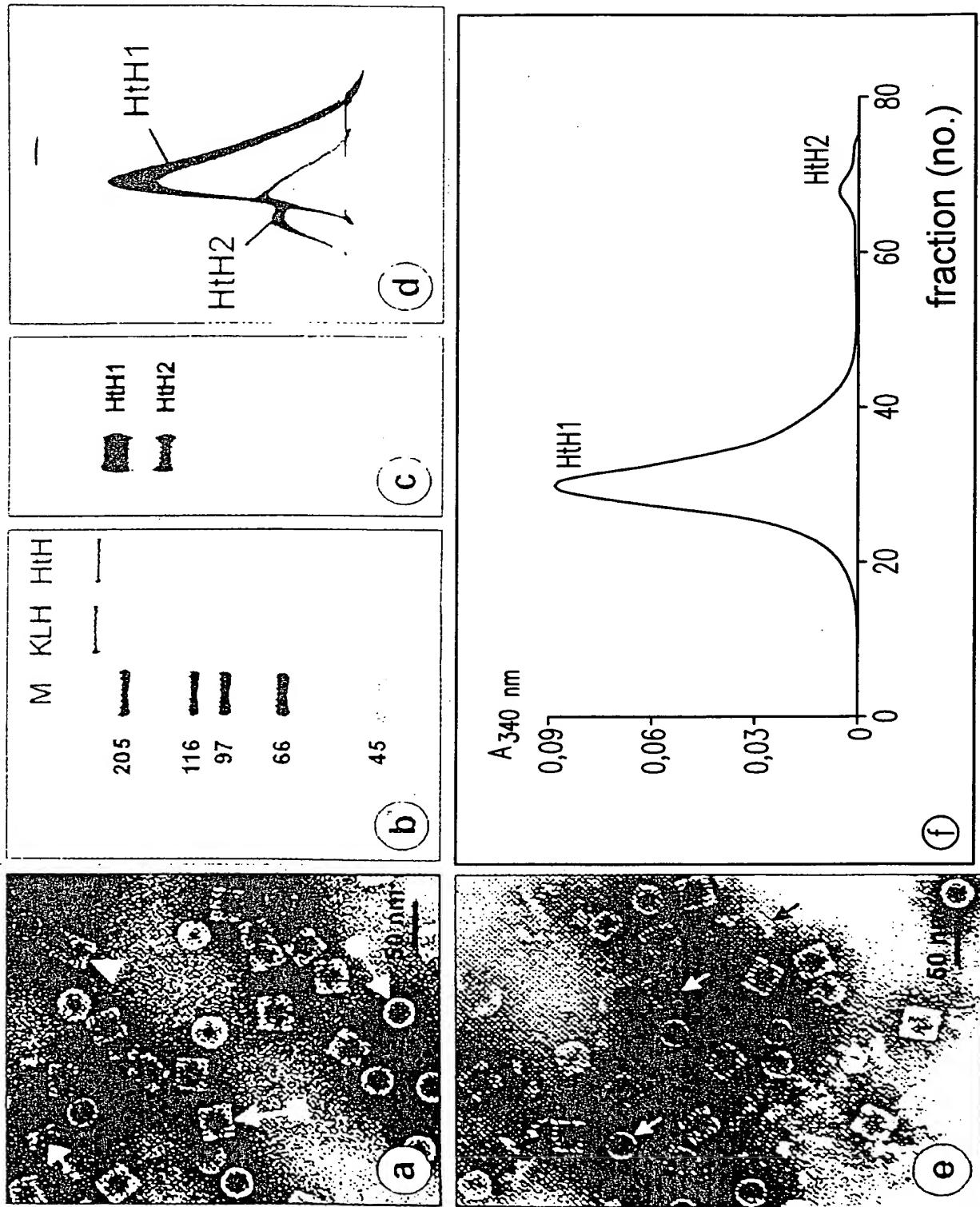


Fig. 1g-m

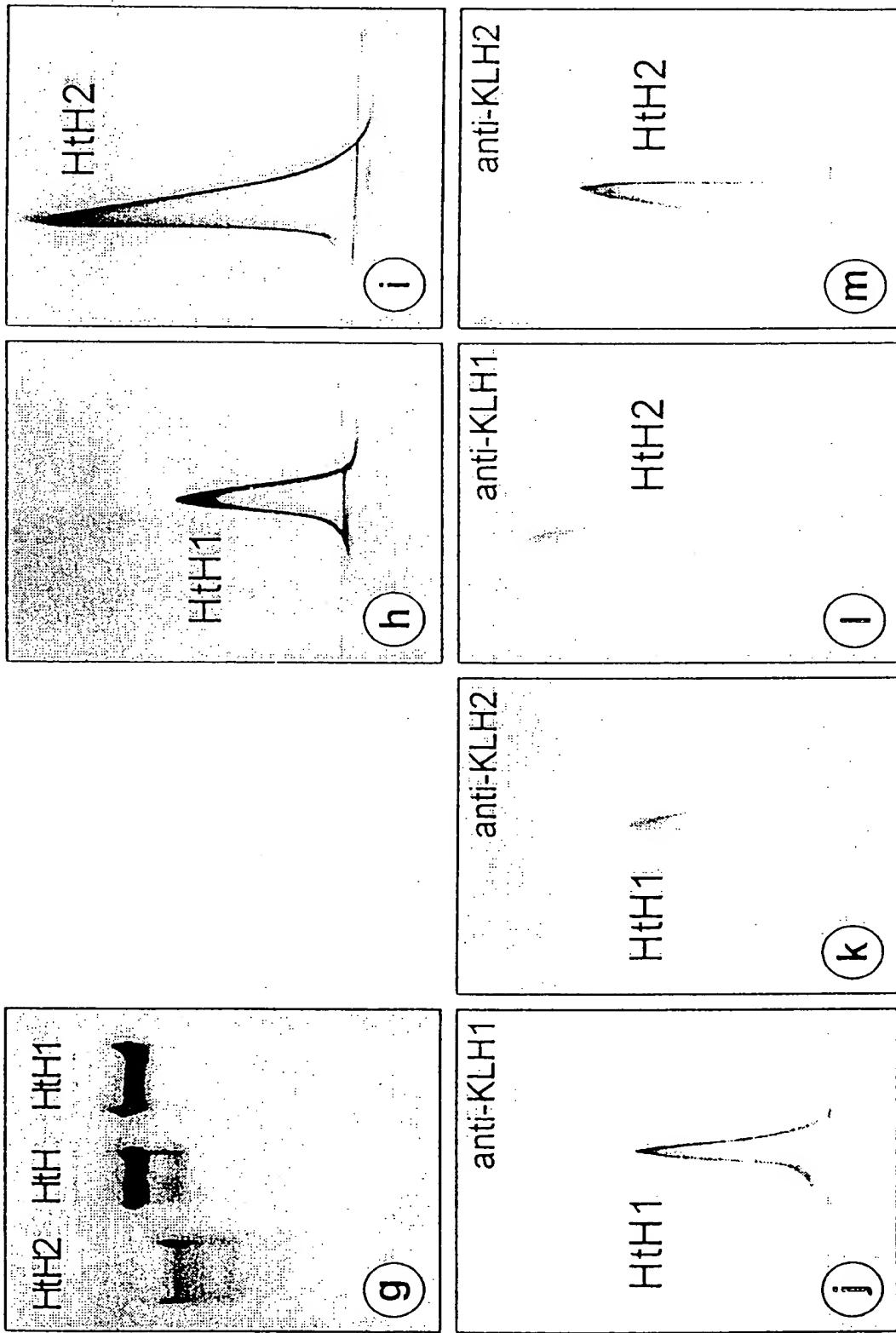


Fig. 2a-h

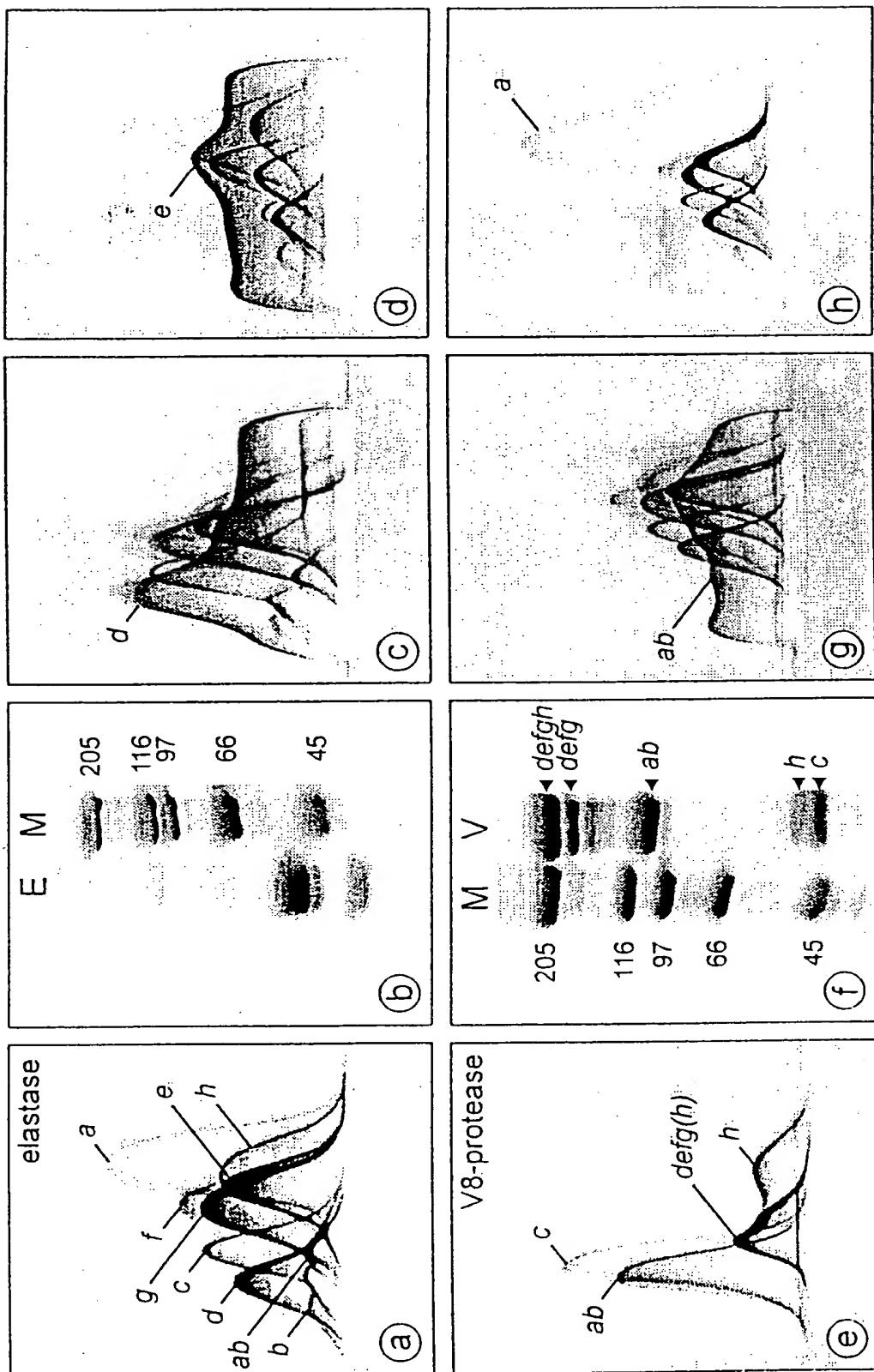


Fig. 2i-p

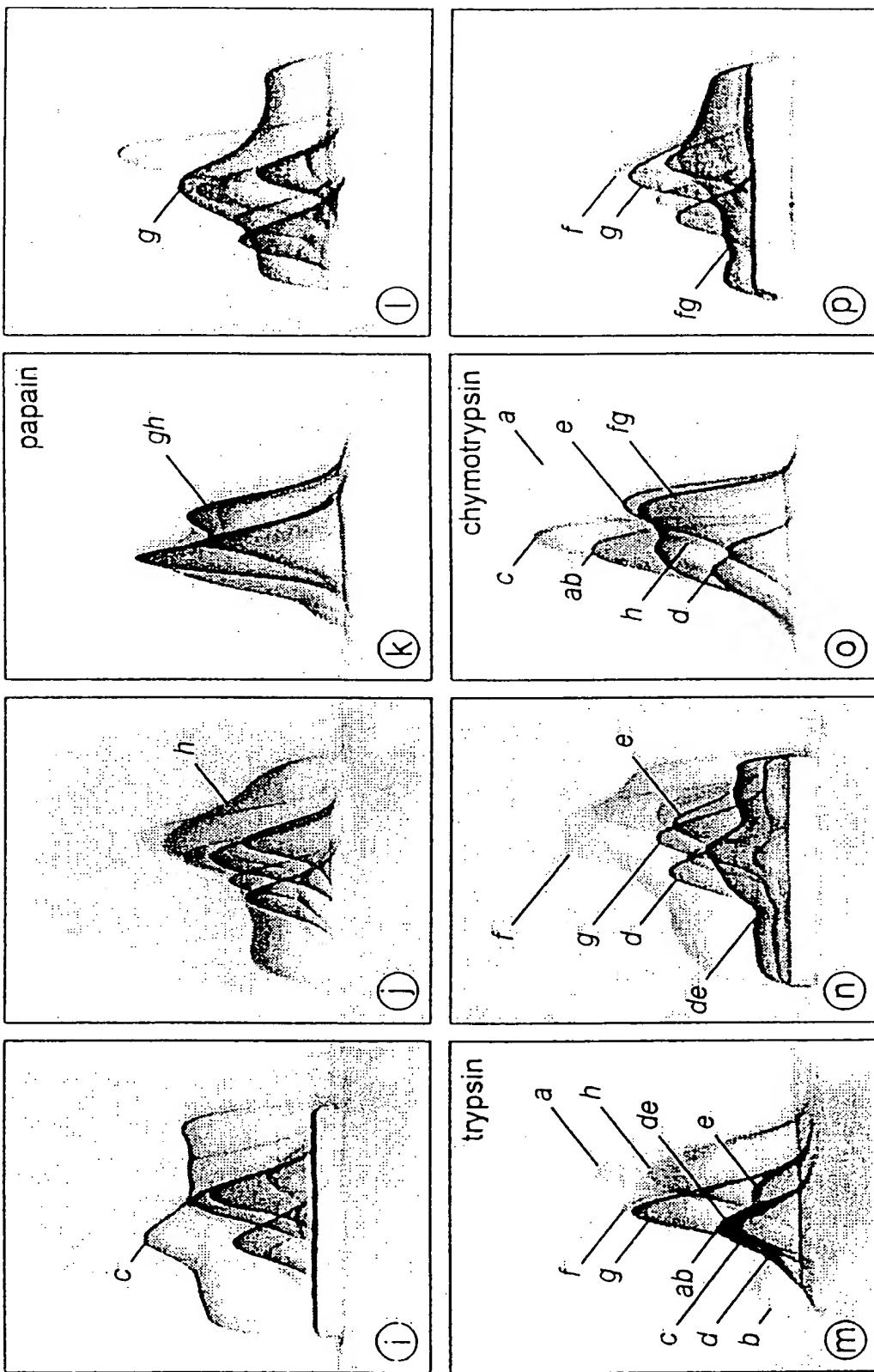


Fig. 3a-c

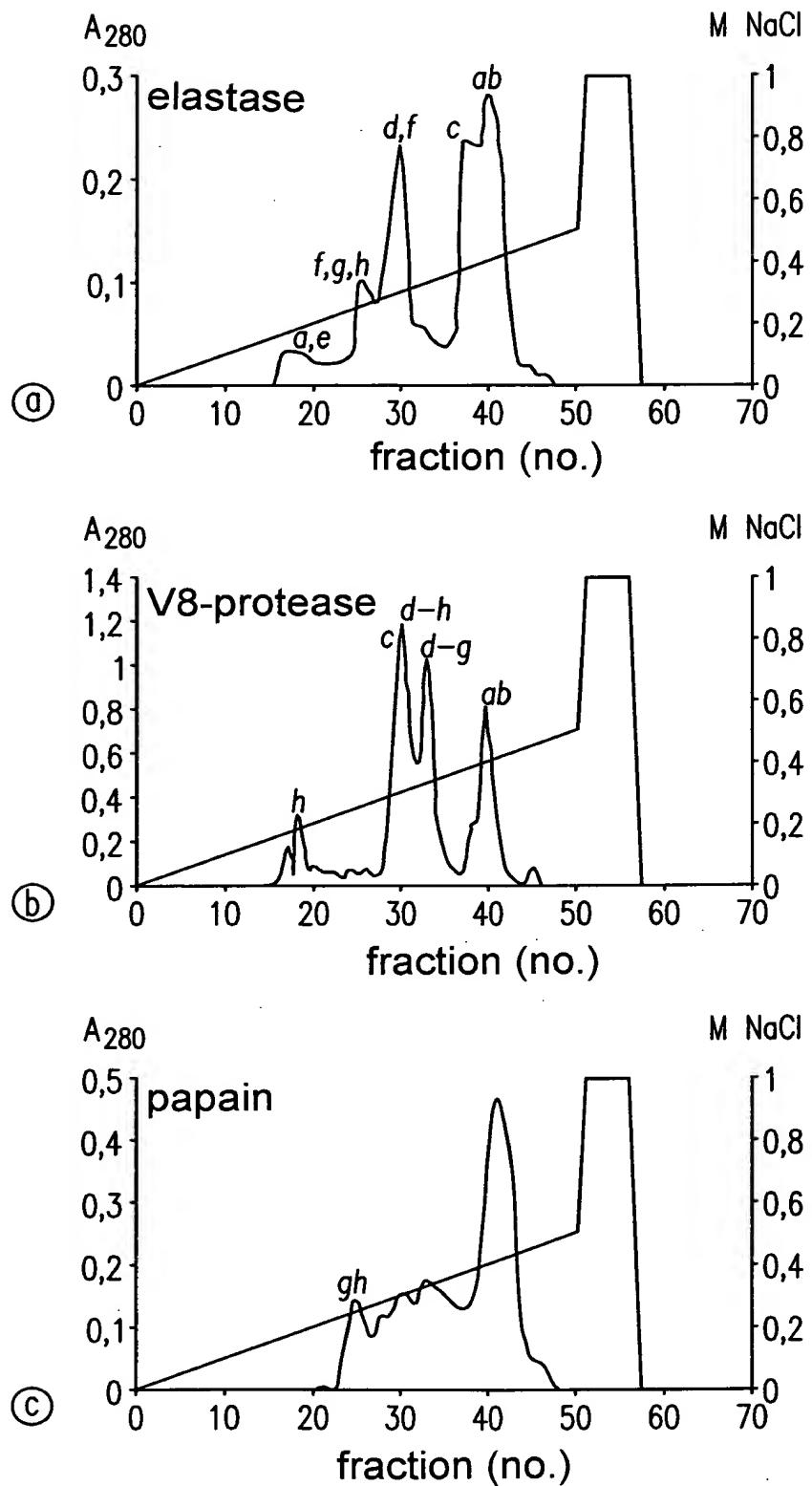
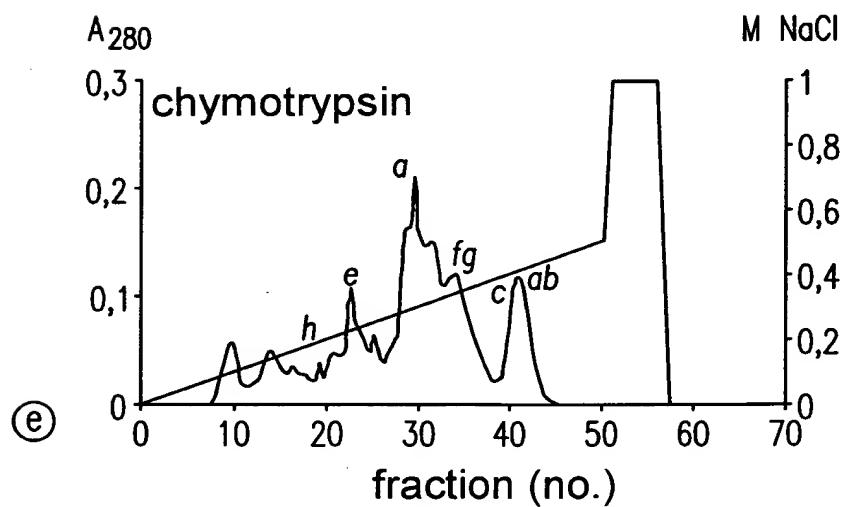
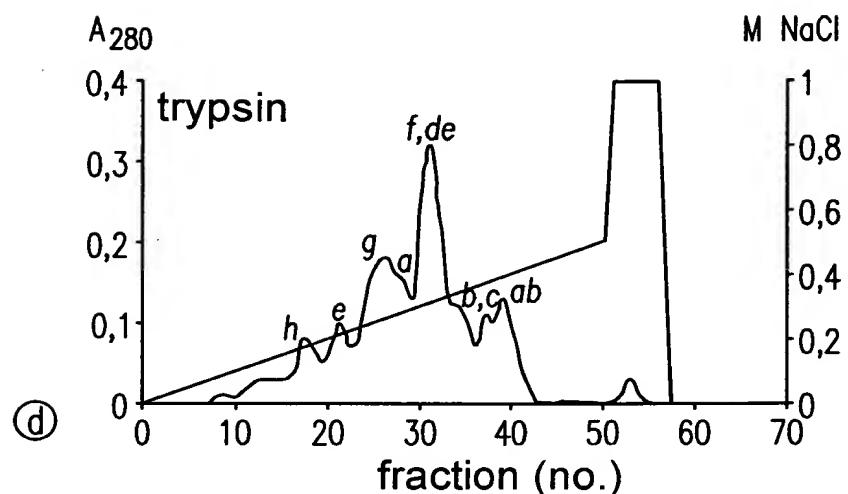


Fig. 3d-e



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Figure 4

HtH1 cDNA sequence and intron structure

Domain a

GGCTTGTTCAGTTCTACTCGTCGCCCTTGTGGCGGGGGCTGGAGCAGACAAACGTCGTCA
AAAGGACGTGAGTCACCTCACGGATGACGAGGTGCAAGCTCTCCACGGGCCCTCCATGAC
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CGTCGTGTGACTACAAGGGACGGAAGATCGCCTGCTGTGTCCACGGTATGCCAGTTCCC
CTTCTGGCACAGGGCATATGTCGTCCAAGCCGAGCAGGGCACTGTTGTC
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GATAGCGTTCATCAATAAGAAGACTGCGCGAGCTGTAGATGATCGCCTATTGAGAAGGTG
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TCTGTAATTTGAAATCCAGTTCGAGTTGGCTCATAATGCTATCCATTACTTGGTTGGCGG
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CGTGGAGATTCTACAGACCCTCTTCTATGATGTA
ACTGAAGCGGTACATCACCTTGGAGT
CCCGCTAAGTGGCACTACTATGTGAAAACAGAACTCTCAGCGTGAATGGCACAGCACT
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Domain b

GTCACTTGACCCACCTGTGCATCATGCCACGATGACGATCTTATTGTTGAAAAAAAT
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TCTGCACATTGAGATTCACTGTTGAGTTAGTCATAATTCTATTGCGTGGATAGGC
TTCCGAAGATTACTCCATGCCACCTGCATTACACAGCCTTGACCCATTCTACCT
CATCATTCAATGTCGATCGTCTATGGCAATCTGGCAAGCTCTCAAATCAGGAGACACA
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AGATTATGAAGTCACTTAAAATTGATGATGACCGAAACGCTCTGATACGGACCTG
ATTCCACACGCAGCAGTTGTTCTGAGCCAGCTCACC

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Intron b/c

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TCGTTGTGTAATGCCACAGCCAGCAATTCCAGATATATAGCGACGGTCTATGAATACTCCA
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TGACTGAAATCTCTCAACGCCGTTAGCAATAATAGGCTCAGTAGTATTCAACCAATTAC
AATCAGTAGAAAATTCTCTATACTATTCTATGTTGCATCCTGATATCCCTATGCAAAAAT
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ATTTCCTTCAG

Domain c

CTACCTTGAGGATGAAAAGCACAGCTTACGAATCAGAAAAATGTCGACAGCTGACTCC
TGAAGAAACAAATGAACTGCGTAAAGCCCTGGAGCTTCTGAAAATGATCATACTGCAGGT
GGATTCAATCAGCTGGCGCTTCCATGGAGAGCCTAAATGGTGCCTAATCCTGAAGCGG
AGCACAAGGTTGCATGCTGTGTTCATGGCATGGCTGTTCCCTCATTGGCACAGGCTTCT
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Intron c/d

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Domain d

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Intron d/e

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Domain e

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Intron e/f

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Domain f(2)

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Intron f/g

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Intron g(2)

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TAGTCAAACAGGTAGGCAGTCCATAAGCATAGTTTAAGCATTAGATCATCTATAATT
AACACATGGTTAGCCGCTATGTTAGTTAACATGAGTTAGAAGTGTATATTG
GAAGGGAAAGTGAAGTAAATCCTTATTCTTACTACCATTAAAGATTCCAATGACTCC
ATTCAACTCCTAACATTACACTGCTCTTCAACAG

Domain g(2)

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GGTGGCCACAGCCCCCTACGGAATGTCCACTCTGACATTCACTGCCTACGATCCTCTTCT
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Intron g/h

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CGTTTAATGGACATGCCCTGTTAATGAAAGGGTAAGTACATGTTATGGGGATGGGATG
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Domain h

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3' UTR

TTCACAG

Intron UTR

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AGCCAAGACTCAACGGGCAGCCGGGTGGGGGATTGGTGGCTGTTCAGACCAGGG
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3' UTR

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ATGGATGTATGTAATGTGGCCGGCAATAAGTATATGTATAAAAAAAAAAAAAAA
A

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Figure 5

Derived primary structure of HtH1

Signal peptide

LVQFLLVALVAGAGA

Domain a

DNVVRKDVSHTDDEVQALHGALHDVTASTGPLSFEDITSYHAAPASCDYKGRKIAACVHG
MPSFPFWHRAYVVQAERALLSKRKTVGMPYWDWTQTLTHLPSLVTEPIYIDSKGKQATNY
WYRGEIAFINKKTAARAVDDRLFEKVEPGHYTHLMETVLDALEQDEFCKFEIQFELAHNAIH
YLVGGKFEYSMSNLEYTSYDPIFFLHHSNVDRLFAIWQRLQELRGKNPNAMDCAHELHQ
LQPFNRDSNPVQLTKDHSTPADLFDYKQLGYSYDSLNLNGMTPPEQLKTELDERHSKERAFA
SFRLSGFCCSANVVVYACVPDDDRSDDYCEKAGDFFILGGQSEMPWRFYRPFYDVTEAV
HHLGVPPLSGHYYVKTELFVNGTALSPDLPQPTVAYRPGK

Domain b

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PRPDAKVRFACCMHGMAFPWHRLFVTQVEDALVRRGSPIGVPYWDWTKPMTHLPDLASN
ETYVDPYGHHTHNPFFNANISFEEGHHHTSRMIDSKLFAPVAFGEHSHLFDGILYAFEQED
FCDFEIQFELVHNSIHAWIGGSEDSMATHYAFDPFIFYLHHSNVDRLWAIWQALQIRRH
KPYQAHCAQSVEQLPMKPFAPSPLNNEKTHSHSVPTDIYDYEEVLHYSYDDLTFGGMNL
EEIEEAIHLRQQHERVFAGFLLAGIGTSALVDIFINKPGNQPLKAGDIAILGGAKEEMPWAF
DRLYKVEITDSLKTLSDLVDGDYEVTFKIHDHMGNALTDLIPHAADVSEPAH

Domain c

PTFEDEKHSLRIRKNVDSLTPPEETNELRKAELLENDHTAGGFNQLGAFHGEPKWCPNPEA
EHKVACCVHGMASFPHWHRLALQAEANALRKHGYSALPYWDWTRPLSQLPDLVSHEQYTD
PSDHVVKHNPWFNGHIDTVNQDTTRSVREDLYQQPEFGHFTDIAQQVLLALEQDDFCSFEV
QYEISHNFIHALVGGTDAYGMASLRYTAYDPFIFYLHHSNTDRIWAIWQSLQKYRGKPNTA
NCAIESMRRPLQPFGLSSAINPDRITREHAI PFDVFNYRDNLHYVYDTLEFNGLSISQLDR
ELEKIKSHERVFAGFLLSGIKKSALVKFEVCTPPDNCHKAGEFYLLGDENEMAWAYDRLFK
YDITQVLEANHLHFYDHLFIRYEVFDLKGVSLGTDLFHTANVVHDSGT

Domain d

GTRDRDNYVEEVGTGASHIRKNLNDLNTGEMESLRAAFLHIQDDGTYESIAQYHGKPGKCQL
NDHNIACCVHGMPTFPQWHLRYVVQVENALLNRGSGVAVPYWEWTAPIDHLPHFIDDATYF
NSRQQRYDPNPFFRGKVTFENAVTRDPQAGLFNSDYMENVLLALEQENYCDFEIQFELV
HNALHSMLGGKGQYSMSSLDYSAFDPVFFLHHANTDRLWAIWQELQRFREL PYEEANCAIN
LMHQPLKPFSDPHENHDNVLKYSKPQDGFDYQNHFGYKYDNLEFHHSIPSLSATLKQRR
NHDRVFAFGFLLHNIGTSADITIYICLPDGRGNDSCHEAGTFYIILGGETEMPFIFDRLYKE
EITKPLQQLGVKLHGGVFELELEIKAYNGSYLDPHTFDPTII FEPGT

Domain e

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DTHILDHDHEEEILVRKNIIDLSPRERVSLVKALQRMKNDRSADGYQAIASFHALPPLCPN
PSAAHRYACCVHGMATFPQWHRLYTVQVQDALRRHGS LGV I PYWDWTKPVNELPELSSAT
FYHPIRNINISNPFLGADIEFEGPGVHTERHINTERLFHSGDHDGYHNWFFETVLFALEQE
DYCDFEIQFEIAHNGIHTWIGGS AVYGMGHLHYASYDPIFYI HHSQTDRIWAIWQELQKYL
GLSGSEANCAIEHMRTPLKPFSGPPYNLNSHTQEY SKPEDT FDYKKFGYRYDSLELEGRS
ISRIDE LIQQRQEKDRTFAGFLLKGFGTSASVSLQVCRVDHTCKDAGYFTI LGGS AEMPWA
FDRLYKYDITKTLHDMNLRHEDTFSIDVTITSYNGTVLSGDLIQTPSIIFVPGR

Domain f

HKLNSRKHTPNRVRHELSSLSSRDIASLKAALTSLQHDNGTDGYQAIAAFHGVPACHEPS
GREIACCIHGMATFPWHRLYTLQLEQALRRHGS VAVPYWDWTKPITELPHILTDGEYYD
VWQNAVLANPFARGYVKIKDAFTVRNVQESL FKMS SFGKHSLLFDQALLALEQTDYCDFEV
QFEVMHNTIHYLVGGRQTYAFSSLEYSSYDPIFFIHHSFVDKIWAVWQELQSRRHLOFRTA
DCAVGLMGQAMRPFNKDFNHNSFTKKHAPVNTVFDYEDLGNYDNLEISGLNLNEIEALIA
KRKSHARVFAGFLFGLGTSADIHLEICKTSENCHAGVIFILGGS AEMHWAYNRLYKYDI
TEALQEFDINPEDVFHADEPFFRLS VAVNGTVIPSSH LHQPTIIYEPGE

Domain g

DHHDDHQSGSIAGSGVRKVNTLTKAETDNLREALWGVMDHGPNQFQAIAAFHGKPALCP
MPDGHNYS C CTHGMATFPWHRLYTKQMEDAMRAHGS HVGLPYWDWTAFTLPTLVTDTD
NNPFQHGHIDYLNVSTTRS PRDMLFNDPEHGS E FFYRQVLLALEQTD FCKFEVQFEITHN
AIHSWTGGHSPYGMSTLDFTAYDPLF WLHH SNTDRIWAVWQALQEYRGLPYNHANCEIQAM
KTPLRPF SDDINHNPVTKANAKPLDVFEYNRLSFQYDNLIFHGYSIPELDRVLEERKEEDR
I FFAFLLSGIKRSADVVFDICQPEHECVFAGTFAILGGELEMPWSFDRLFYDITKVMKQL
HLRHDS DFTFRVKIVGTDDHELPDSVKAP TIEFEPG

Domain h

VHRRGGNHEDEHHDDRLADVLIRKEVDFLSLQEANAIKDALYKLQNDDSKGGFEAIAGYHGY
PNMC PERGTDKYPCCVHGM PVFPHWRLHTI QMERALKNHGSPMGIPYWDWTKKMSSLPSF
FGDSSNNNPFYKYYIRGVQHETTRDVNQRLFNQTKFGEFDYLYLTLQVLEENSYCDFEVQ
YEILHNAVHSWLG GTGQYSMSTLEYSAFDPV FMIHHSSLDRIWILWQKLQKIRMKPYYALD
CAGDRLMKDPLHPFNYETVNEDEFTRINSFPSILFDHYRFNYEYDNMRIRGQDIHELEEV
QELRNKDRIFAGFVLSGLRISATVKVFIHSKNTSHEEYAGEFAVLGGEKEMPWAYERMLK
LDISDAVHKLHV KDEDI RFRVVVTAYNGDVVTTRLSQPFIVHRAHVAHDILVI PVGAGHD
LPPKVVVKSGTKVEFTPIDSSVNKAMVELGSYTAMAKCIVPPFSYHGFELDKVYSVDHGDY
YIAAGTHALCEQNLRLHIHVEHE

15/29

Figure 6

HtH2 cDNA sequence and intron structure

Domain b

CACAGACTGTCGTCACCCAGGTGGAAGATGCTCTGATCAGGCGAGGATGCCTATAGGGG
 TCCCCTACTGGGACTGGACTCAGCCTATGGCGCATCTCCAGGACTTGCAAGACAACGCCAC
 CTATAGAGATCCCATCAGCGGGGACAGCAGACACAACCCCTTCCACGATGTTGAAGTTGCC
 TTTGAAAATGGACGTACAGAACGTACCCAGATAGTAGATTGTTGAACAACCTTATTG
 GCAAACATACGCGTCTTCGACAGTATAGTCTATGCTTTGAGCAGGAGGACTTCTGCGA
 TTTGAAAGTTCAATTGAGATGACCCATAATAATATTACGCGCTGGATTGGTGGCGCGAG
 AAGTATTCCATGTCCTCTACACTACACAGCCTCGACCCATCTTCTACCTTCGTCACT
 CCAACACTGACCGGCTCTGGGCAATTGGCAAGCGTGCAGATACGAAGAACAGGCCTTA
 CAAGGCTCATTGTGCTTGGCTGAGGAACGCCAGCCTCTCAAACCTTCCAGTTCC
 CCACTGAACAACAACGAAAAACTACGAAAACCTCGGTGCCACCAACGTTACGACTACG
 AAGGAGTCCTGGCTATACTTATGATGACCTCAACTTCGGGGCATGGACCTGGGTAGCT
 TGAGGAATACATCCAGAGGCAGAGACAGAGAGACAGGACCTTGCTGGTTCTTCTGTCA
 CATATTGGTACATCAGCGAATGTTGAAATCATTATAGACCATGGACTCTTCATACCTCCG
 TGGGACGTTGCTTGGCTGAGGAAGAGATGAAATGGGGATTGACCGTTGTA
 CAAATATGAGATTACAGATGAACTGAGGCAACTTAATCTCCGTGCTGATGATGTTTCAGC
 ATCTCTGTTAAAGTAACTGATGTTGATGGCAGTGAGCTGTCCTCTGAACTCATCCCATCTG
 CTGCTATCATCTCGAACGAAGCCATA

Intron b/c

GTAAGTAGCTACCTGTTATTCAATTTCGCTTGCCTAACATTCAATTGAGCTTGAAA
 TTCAATAATTGTTGCTGGCTGAAACCAATTGAACTCTTCTTCTCAGGTG
 AACTCAAATAATACTCAATTGTTATGCACGCGGGTAGGGCATACTACTATCCAC
 ATCGGTCACTCTCAAAATGCAAACAAATTGCTTATTTCGTTGGACAAGCAAACCCCT
 TCCTGTAATCTGCCTTGGCATCCACTGGAATTATGTTGACTGGTAATTGATACTGGCT
 CTCTCTTGCTAGAGTTAATATCTATAGTTGAAATCTTATGATTGCTATTAT
 TTCGACAGCATGCTATAGACACCCCTAGACTATTGATAGCCACTGTTGATTGTTCCATT
 TATTATTATAACAGAACATGGCTTGTATTACCTTCCAG

Domain c

TTGACCATCAGGACCCGCATCATGACACAAATCATTAGGAAAAATGTTGATAATCTTACACC
 CGAGGAAATTAAATTCTCTGAGGCGGGCAATGGCAGACCTCAATCAGACAAACCGCCGGT
 GGATTCCAGCAAATTGCTGCTTTCACGGGAACCAAATGGTGCCAAGTCCGATGCTG
 AGAAGAAGTTCTCTGCTGTCCATGGAATGGCTGTCTCCCTCACTGGCACAGACTCCT
 GACCGTGCAAGGCGAGAATGCCCTGAGAAAGCATGGATGTCTGGAGCTCTCCCTACTGG
 GACTGGACTCGGCCCTGTCACCTACCTGATTGGTTGGTAAGTAGCAGAACTACAC
 CGATGCCATATTCCACCGTGGAAAGCCCAGAACCCCTGGTACAGCGGCCATATTGATA
 CGTGGTGTGACACAACAAGAGCGTCCGTCAAGAACTGTATGAAGCTCTGGATTGGCCAT
 TATACTGGGTCGCTAAGCAAGTGCCTCTGGCTTGGAGCAGGATGACTCTGTGATTG
 AAGTCCAGTTGAGATAGCTCACAAATTCTACGCTCTGTCGGCGGAAGCGAGCCATA
 TGGTATGGCGTCACTCCGTTACACTACTATGATCCATTCTACCTCCATCATTCTAAC
 ACTGACAGACTCTGGGCTATATGGCAGGCTCACAAAGTACAGGGGCAAACCTACAATT
 CCGCCAACGTGCGCCATTGCTTCTATGAGAAAACCCCTACAACCCCTTGGTCTGACTGATGA
 GATCAACCCGGATGATGAGACAAGACAGCATGCTGTTCTTCAGTGTCTTGATTACAAG
 AACAACTTCAATTATGAATATGACACCCCTGACTTCAACGGACTATCAATCTCCAGCTGG
 ACCGTGAACGTGTCACGGAGAAAGTCTCATGACAGAGTATTGCGGATTGCTGCATGG

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TATTCAAGCAGTCTGCACTAGTTAAATTCTTGTCTGCAAATCAGATGATGACTGTGACCAC
TATGCTGGTGAATTCTACATCCTTGGTATGAAAGCTGAAATGCCATGGGCTATGATCGC
TTTACAAATATGAGATCACTGAGCAGCTCAATGCCCTGGATCTACACATCGGAGATAGATT
CTTCATCAGATACGAAGCGTTGATCTTCAATGGTACAAGTCTGGAAAGCAACATCTTCCCC
AAACCTCTGTACATGACGAAGGGCAG

Intron c/d

GTGAGAACATTGATAATAGTTCAAATgAAGTATATCCGATTCAAGCTGTCGATACAAGATg
AGATAACATAATCACAATGTTGTATTAGATATCTCTCTTAATTAAATGCCGCTTTATCAA
TATTGAGCAATCCTCAGCAACATACACCAGCAAATGTTCATCAACAGACTATATTATT
TAATCTTTAAAATCCTTCTGTTATAAAACTTAAAGTATCGAATTCTTGAATG
CGTCTCTGCAGCATAGTTAAGTGTGTTCTGTCAAG

Domain d

GTCACCATTAGGCTGACGAGTACGACGAAGTTGTAAGTGCAGGCCACATCAGAAAGAA
TTTAAAAGATCTGTCAAAGGGAGAAGTAGAGAGCCTAAGGTCGCTTCCCTGCAACTTCAG
AACGACGGAGTCTATGAGAATATTGCCAAGTTCCACGGCAAGCCTGGGTTGTGATGATA
ACGGTCGCAAGGTGCTGCTGTTGTCCATGGAATGCCACCTCCCCAGTGGCACAGGCT
CTATGTCCTCCAGGTGGAGAATGCTTGCTGGAGAGAGGATCTGCCGTCTGTGCCATAC
TGGGACTGGACTGAAACATTACAGAGCTGCCATCTTGATTGCTGAGGCTACCTATTCA
ATTCCCGTCAACAAACGTTGACCCATACTCTTCTCAGAGGTTAAACAGGTTACTACCAAAAC
TGCTGTTACAACACGTGATCCCCAGCCTGAGCTGTACGTTAACAGGTTACTACCAAAAC
GTCATGTTGGTTTTGAACAGGACAACACTACTGCGACTTCGAGATAACAGTTGAGATGGTC
ACAATGTTCTCATGCTTGGCTGGTGGAAAGAGCTACTTATTCTATTCTCTCTGATTA
TTCTGCATTGACCCCTGTGTTTCTCACCATGCGAACACAGATAAGATTGTGGGCCATC
TGGCAGGAGCTGCGAGGGTACAGGAAGAAGCCATACAATGAAGCGGATTGTGCCATTAACC
TAATGCGAAACCTCTACATCCCTGACAAACAGTGTACATCATGATCCTGTAACCTT
TAAATACTCAAAACCCACTGATGGCTTGACTACCAGAACAACTTTGGATACAAGTATGAC
AACCTTGAGTTCAATCATTCACTGTTGACTTCCAGGCTGAAAGAAATCATTGTTATTAGACAAAC
GTCAAGATCGTGTGTTGCGAGGATTCTCCTCACAACATTGGGACATCCGCAACTGTTGA
GATATTCCCTGTGTCCTACCAACCCAGCGGTGAGCAAAACTGTGAAAACAAAGCCGGAAACA
TTTGCCTGACTCGGAGGGAGAACAGAGATGGCTTCTATTGACAGACTCTACAGGTTG
ACATCAGTGAACACTGAGGGACCTCGGCATACAGCTGGACAGCCATGACTTGTGACCTCAG
CATCAAGATTCAAGGAGTAAATGGATCCTACCTGATCCACACATCCTGCCAGAGCCATCC
TTGATTTGTGCCCTGGTTCAAGT

Intron d/e

AAGAAAGTTCACTGTCTAAATCTTTTATGATAGAGGGTAGAGAAGTGGAGACAATGT
GACAATATATTGAATAAAGTTGTTAAAATTATAACTCTCATAGTTCAATTATGCTGA
AGCTGTAGCCATCTATAACTGTGTAACATGAAATGTTAACAGACATTAACCTAAATACTTCAG
CTGATAACAAACAAATGTTAACATACGTCATGTAACATTCTTATCTTAGGTTATA
GCATAAACACTTCAGAGATAACAGTGACGAAACCTCTATTAAATATTCAAGGT

Domain e

TCTTCCTGCGTCTGATGGCATTCAAGATGACATCCTGAGAAAAGAAGTGAACAGCC
TGACAACCAGGGAGACTGCATCTGATCCATGCTCTGAAAAGTATGCAAGGAAGACCATT
ACCTGACGGGTTCCAAGCCATTGCCCTTTCCATGCTCTGCCACCACTCTGCCCTTCACCA
TCTGCAGCTCACCGTTATGCTGCTGTGTCACGGCATGGCTACATTCCCCAGTGGCACA
GATTGTACACTGTACAGTTCCAGGATGCACTGAGGAGACATGGAGCTACGGTAGGTGTACC
GTATTGGGATTGGCTGCGACCGCAGTCTCACCTACCAAGAGCTTGTACCATGGAGACATAC

REPLACEMENT SHEET NO. 17 }

17/29

CATGATATTTGGAGTAACAGAGATTCCCCAACCTTCTACCAAGCCAATATTGAGTTG
AAGGAGAAAACATTACAACAGAGAGAGAAGTCATTGCAGACAAACTTTGTCAAAGGTGG
ACACGTTTTGATAAACTGGTTCTCAAACAAAGCCATCCTAGCGCTGAGCAGGAAAAC
TGTGACTTTGAGATTCAAGTTGAAATTCTTCACAACGGCGTCACACGTGGTCCGGAGGCA
GTCGTACCTACTCTATCGGACATCTCATTACGCATTCTACGACCCTCTTCTACCTCA
CCATTTCAGACAGACCGTATTGgGCAACTGGCAAGAACTCCAGGAACAGAGAGGGCTC
TCGGGTGATGAGGCTCACTGTGCTCTGAGCAAATGAGAGAACATTGAAGCCTTCAGCT
TCGGCGCTCCTTATAACTGGAATCAGCTCACACAGGATTCTCCCGACCCGAGGACACCTT
CGACTACAGGAAGTTGGTTATGAATATGACAATTAGAATTCTGGAAATGTCAGTTGCT
GAACTGGATCAATACATTATTGAACATCAAGAAAATGATAGAGTATTGCTGGTCTGT
TGAGTGGATTGGAGGTTCCGCATCAGTTAATTCCAGGTTGTAGAGCTGATTCCACATG
TCAGGATGCTGGGACTTCACCCTTGGCAGTGCTGAGATGGCGTGGGATTTGAC
AGGCTTACAAATATGACATTACTGAAACTCTGGAGAAAATGCACCTCGATATGATGATG
ACTTCACAATCTCTGTCAGTCTGACCGCCAACACGGAACTGTCCTGAGCAGCAGTCTAA
CCCAACACCGAGTGTCAATTCCAGCGGGGACATC

Intron e/f

AAGTAGTAAACTGCTCAGATTGTTCTAAATTACTCCACTATTAAGTAAAAGTACTAGT
AATTCAATAGTACTGTTCACAGAGAAATGTAACACAATAGACACAGAGTCCTTTGTTAA
ACGCCTTGGCTGGTAAGTCTGAGGTTGGTACTGATGAAAGCTAAAATATTTG
ACAG

Domain f (1)

GTGACATAAATACCAAGGAGCATGTCACCGAACCGTGGTCCGCGTGAGCTGAGCGATCTGTC
TGCAGGGACCTGCTAGTCTCAAGTCTGCTCTGCGAGACCTACAGGAGGATGATGGCCCC
AACGGATACCAGGCTCTGCAGCCTCATGGCTACCAGCAGGCTGCCATGATAGCCGGG
GAAATGAGAT

Intron f

ATATTTAAAGTATTTATCTACGCATGACCCCTGACCCATTATTTTTAATCCTATGAT
GAAACATTACTTAGCTGGCTTGAGCCCCAGGCAAAATGCACTGAAAAATACACTGA
CAGAGGATTAGGCATTCTGGGAGTACTGTATAGTTAGTGCATACATATTAGCGTTCCCT
CACTAAAACGAATCTCTGAATGCTATCAATTAAAGATCATGATGCTTGATTGTTCTACT
GTATTTAAATGGTTAAGATTGCAATTACAATATACACAAACACGTTCTGCATCTC
GGAGAATGCAATCTTCGTTGTACGCGTCTGTTCATATTTTATGCATGTTGAC
TACTTAGCGTCCAATAATCCATTACAAAATCACACAAACAAACGATTTAGGAATGTGA
CTGTAGCTGCAACGAATATACCTGATCCTTCTTGTCCAGAT

Domain f (2)

CGCATGTTGCATTCACGGGATGCCGACCTTCCCCAGTGGCACAGACTGTACACCCCTGCAG
TTGGAGATGGCTCTGAGGAGACATGGATCATCTGTCGCCATCCCTACTGGGACTGGACAA
AGCCTATCTCGAACTCCCTCGCTTCAACAGCCCTGAGTATTATGACCCATGGCATGA
TGCTGTGGTAAACAAACCCATTCTCAAAGGTTGTCAAATTGCAAATACACTACAGTA
AGAGACCCACAGGAGATGCTGTTCCAGCTTGTGAACATGGAGAGTCATCCTCTATGAGC
AAACTCTTCTGCTCTGAGCAAACCGACTACTGTGATTTGAGGTACAGTTGAGGTCT
CCATAACGTGATCCACTACCTGTTGGACGTCAACCTACGCATTGCTTCTGCAT
TATGCCTCTACGACCCATTCTTATACACCATCCTTGTGGATAAGATGTGGGTAG
TATGGCAAGCTCTCAAAAGAGGAGGAAACTCCATACAAGCAGCTGACTGTgCTGTC
CCTAATGACTAAACCAATGAGGCCATTGACTCCGATATGAATCAGAACCCATTCAAAAG
ATGCACGCAGTTCCAACACACTCTATGACTACGAGACACTGTAACAGCTACGATAATC
TCGAAATAGGTGGCAGGAATCTGACCCAGCTCAGGCTGAAATTGACAGAAGCAGAACCA

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CGATCGCGTTTGTGGATTCTGCTTGCAGAATCGGAACCTCTGCTGATGTCAGGTT
TGGATTTGAGAAATGAAATGACTGCCACAGGGGTGGAAATAATTTCATCTTAGGTGGAG
CCAAGGAAATGCCATGGTCATTTGACAGAAACTCAAGTTGATATCACCCATGTACTCGA
GAATGCTGGCATTAGCCCAGAGGACGTGTTGATGCTGAGGAGCCATTATATCAAGGTT
GAGATCCATGCTGTTAACAGACCATGATACCGTCGTGATCCCAGCCCCAACTATCA
TCTATTCTCCTGGGGAAAG

Intron f/g

GTGAGAGAACCACTAATAGCTACTGTCTACAAAGAACATGTGTTCATTTAAAGACCTGACTGT
AGGCCGATGGCTGCTGTCATCTCCTCCGCCTCCCTGTCCTCCTCCGAAGGGGTCA
GCTTCAGGTTCTTGCCAAATATGCCAACGACCTCCCTGAGCAGGAGTATATACGTA
AGGGAAAGCAAGTATGGACCATCGCGCGCATGTAGAGATAACATGATCAGCTGCTGCTGT
TCCACTCCTGTCAGACAATGAGATAAACATGAATAACAGTATTACTCAGCAGCGTTCCAATT
TTCAACCCTCGTATTTATTAAAAAAAGGAATTTAATATATTTCTCCTGTTGAAATA
TTTAGTAACTGTTAATCGATATAGAGTGGAGTAGTGACGCTTATTTCGGtTCATTCTCG
AAACAAAAATATAATAGTCCACTGAACCTCTTAAATTGTTTACAACCTCAACTGCCA
CAGACGTAATCCCTCACGTATTGAGCTGACAACGTGTTGAATTGAGTGTGTTCCGAAT
TCTAAATAAGCATGTATATATTACGTCTCATGCAAGTAATAATATGTTAACTGATGACGT
CACTGGTGACCACTGATTAGTCCCTTGTCAGCTGAATATCCCCTGCAATCCCCACATGGGT
GGTGGGGAAAGCCAGGTTCCCTGTCACGCTGAATATCCCCTGCAATCCCCACATGGGT
ACAAAGTGTGATGCCATTCTGGTGTCCCCACCGTGATATTGCTGGAATAAGTGGCTTA
ATACCATATACACTCACTCTATTGTCACACTACTGCCACCGGCTCACACCTCTGATGCTTC
TGTTCTATCCAG

Domain g(1)

GTCGCGCTGCTGACAGTGCACACTGCAACATTGCTGGCTCTGGGGTAGGAAAGGACGT
CACGACCCCTCACTGTGTCAGACCGAGAACCTAACAGACAGGCTCTCAAGGTGTCATCGAT
GATACTGGTCCAATGGTTACCAAGCAATAGCATCCTCACGGAAGTCCTCCAATGTGCG
AGATGAACGGCCGCAAGGTTGCCTGTTGCTCACG

Intron g(1)

GTAATTAAATGGATGTGAAGTCAATGTCGAGGGTATAATAAGGATTAAATACTTCAGTCG
TGTAATACTGTATGACATGTGTTGGATGGTAGGTATTACAGGTATAAGGCCAGTGT
GTGTTGGGACGGTTACTTCTGCACTAGTAATAAGCATTGTATTAGCTAGCTTTATCA
TATAACTTTAGTTCAAGGTTGtGGCAATTGAAATCGAAATTTCCTTCATTCAAGGTTA
TCGCACTCGTGTGNAGAATAGTTACTATGCTGTCATTGAGAATAACACTATAGTAATAAG
CATATCATACAGTAAGAATAACACTATAGTAATAAGTATATCATNCAGTAAGAATGTCA
TGTATGATAAAATAGGTTATCACACTCGTGTGTTAGAATGGTTACTATCCCAGGAATAAC
CACTATGTATTACATGTATATTGGGCAGTGTAAAGTAGTAGCATTGTATATTAAATCACT
ATCGTGCTCAAAACACCAGGATATATGGGGTATACAGTGGCAGTGTAAAGTAGCAACATT
GTATATTAAATCAGTATATCGTACTCAAAACACCAGGATTATGGGGTATACAGTGGCAG
TGTAAGTAGTACGATTGTATATTAAATCAGTATATCGTACTCAAAACACCAGGATATAAT
TCAGTATATCGTGTGTTCAAAACACCAGGATATAATTCACTGTTCAAAACACCA
GGATATATGGGATATACTGCGGGTTGCATACAACCTCCACCCCTTACAG

Domain g(2)

GTATGGCCTCTTCCACACTGGCACAGACTGTATGTGAAGCAGATGGAAGATGCCCTGGC
TGACCACGGGTACATATCGGCATCCCTACTGGGACTGGACAACTGCCTCACAGAGTTA
CCCGCCCTTGTACAGACTCCGAGAACAACTCCCTCATGAG

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19/29

Intron g(2)

GTCAGTTAGTCTCCTGTCAGCTAACGATAACCAATTCTATTTGAGAACCCACGATG
ACGAGAAAACAAGCAATATAGATATAGATGCAGTATAGATCAAGTTAATGAATTCTATTGCT
ATATGTTGCTTGTAAATAACTTAAGAAAACGAGAGCATGCACACAAATGAAACAAACAA
TTATGTGTTGATAGGAATATGATATATGTATTGGGGCTGACGTGAGCAGGGTTGAAGG
GACAGTTACATTGTCAGTAACACTGGAGTATTCTTGTATCCACAATATATAGTTCAT
GTGTTCAGCAGTTACAACATTATACATACATTACGTCGAAACATGCTTCTTGTC
CTCTTGTGCCAG

Domain g(3)

GGTCGCATTGATCATCTCGGTGTAACCACGTACGTTCCCCAGAGACATGCTGTTAACG
ACCCAGAGCAAGGATCAGAGTCGTTCTTCTATAGACAAGTCCTGGCTTGGAGCAGAC
TGACTACTGCCAGTTGAAAGTCCAGTTGAGCTGACCCACAACGCCATTCACTCCTGGACA
GGTGGACGTAGCCCTTACGGAATGTCGACCCCTCGAGTTCACAGCCTACGATCCTCTCT
GGCTTCACCACCTCCAACACCGACAGAACTCTGGCTGCTGGCAAGCAGTGCAGAAATACCG
AGGACTCCCATAAACGAAAGCACACTGTGAAATCCAGGTTCTGAAACAGCCCTTGAGGCCA
TTCAACGATGACATCAACCACAATCCAATCACCAAGACTAATGCCAGGCCTATCGATT
TTGATTATGAGAGGTTAACATATCAGTATGACACCCTAGCTTCCATGTTAAGAGCATCCC
TGAACCTGAATGACCTGCTCGAGGAAAGAAAAAGAGAAGAGAGAACATTGCTGCTTCTT
CTTCGTGGAATCGGTTGCACTGCTGATGTCGTTGACATCTGCCGgCCCAATGGTGACT
GTGTCTTGCAGGAACCTTGCTGCTGGAGGGGAGCTGAAATGCCCTGGTCTTCGA
CAGACTGTTCCGCTATGACATCACCAAGACTGAAATCAGCTCCATCTCAGTATGATT
GATTTCAGTTCAGGGTGAAGCTTGTGCCACCAATGGCACTGAGCTTCATCAGACCTTC
TCAAGTCACCAACAATTGAACATGAACATTGG

Intron g/h

GTATGTTATCTTATCATCAAATGTGATCAGATACTGGAGACGTTTATTAACATTGG
TCAGCATTAGTTGATGATTGGTGCAGTGTGACGACAAGGAGTCAGCATTAAACACATT
CAACACATCTTAATCTGATATGAGAAGGGAAATAAATTGATCCAGTATTGATGATTGAAGT
TAGATTAACAGTGAAAGATATACCAGTTTGATAATCGTATAAAACAGTAGCAGAAATTGTA
TCGTGAAAACATAATGTGGGAAGGGCGAACGCCAACGAGATTAGATTACGATCGTGTGCT
AGAATAATTACAATAACCCAGACGTCGGAAATGTGGTGTCTATGCCAATGGTACGATT
AATTGCTAACATGCACGATTACCTATTCAG

Domain h

AGCCCACAGAGGACCAGTTGAAGAAACAGAAAGTCACTCGCCAACATACTGACGGCAATGCA
CACTTCATCGTAAGGAAGTTGATTGCTGTCCTGGATGAAAGCAAACAAACTTGAAGAATG
CCCTTACAAGCTACAGAACGACCACAGTCTAACGGGATACGAAGCAATCTCTGGTTACCA
TGGATACCCCAATCTGTGTCGGAAAGAAGGGCGATGACAAAATACCCCTGCTGCGTCCCCGG
ATGGGCATCTTCTTACTGGCACAGACTCTTGACCACTCAACTGGAAAGAGCTTTGAGC
ACAATGGTGCAGTGCCTGGTCTTACTGGACTGGAAACAAGGACCTGCGTCACTGCC
GGCGTTCTCCGACTCCAGCAACAAACATCCACTTCAAGTACCACTGCCGGTGT
GGTCACGACACCGTCAGAGAGCCAAGTAGTCTTATATAACCAGCCCAATCCATGGTT
ATGATTATCTCTTACCTAGCATTGACCAAGCCTGAGAAGAAAACAATTACTGGACTTTGA
GGTCAGTATGAGATCCTCCACAAACGCCGTCACCTCTGGCTGGAGGATCCCAGAAGTAT
TCCATGTCTACCCGGAGTATTGGCTTGTGACCTGCTTTATGATCCTCACTCGGGTC
TAGACAGACTTGGATCATCTGGCAAGAACTTCAGAAGATCAGGAGAAAGCCCTACAACCT
CGCTAAATGTGCTTATCATATGATGGAAGAGCCACTGGCGCCCTCAGCTATCCATCTATC
AACCAAGGACGAGTTCACCCGTGCCAACTCCAAGCCTTACAGTTTGACAGCCATAAGT

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20/29

TCGGCTACCATTACGATAACCTGAATGTTAGAGGTACAGCATCCAAGAACTCAACACAAAT
CATCAATGACTTGAGAACACAGACAGAACATCTACGCAGGATTGTTGTCAGGCATCGGT
ACGTCTGCTAGTGTCAAGATCTATCTCCGAACAGATGACAATGACGAAGAAGTGGAACTT
TCACTGTCCTGGGAGGGAGAGGGAAATGCCATGGGCCTACGGAGCGAGTTTCAAGTATGA
CATCACAGAGGTTGCAGATAGACTAAAATAAGTTATGGGACACCTTAACCTCCGGA
ACTGGAGATCACATCCTACGAATGGAATCGGTGGTAAACAAGAGCCTACCCAAATCCTT
CATCATCTACAGACCTGCCAATCATGACTACGATGTTCTGTTATCCCAGTANGGAAGAAA
CCTTCACATCCCTCCCAAAGTTGTCGTCAAGAAAGGCACCCGCATCGAGTTCCACCCAGTC
GATGATTCACTACGAGACCAGTTGTTGATCTGGAAGCTACACTGCACACTTCAACTGTG
TGGTACCAACCGTTCACATACCACGGATTGAACTGAACCACGTCTATTCTGTCAAGCCTGG
TGACTACTATGTTACTGGACCCACGGAGAGACCTTGCAGAATGCAGATGTCAAGGATTCA
ATCCATGTTGAGGATGAGTAA

3' UTR

CGCAACAGGT

Intron UTR

GAGATAAGAAACCCCTCTAACAGTAATACGACACCACTACAGCTAAACATGATTGCCA
TCGATTTTCATGTTAGTATACGCTTTCAAGTTCTACATAATTGTTTTCAAATCAA
GTTAGCAAATGAATCTATCACTGGAAAATAGGGTAGGGTAGCCAAGTGGTTAAAGCGGTC
ACTGATCACGCCAAAGACGAGTGTCTAACCTGCATGGGTACAAAAGTGAAGACCATTGCT
GGTGTCTACCGCCGTAATTGTTAGTATTGCTAAAACCTATACACCCATGCGCTG
TAAAAGTGGAAATAATAATCATATTCAACAAAAGCACAAACATTCAATTTCATGAAAG
CCTCTGTTACCTGAAAGACGCAAGAGAACATAGTTCTAACATTATTCAGACATG
GAAATGTCCTGCACGTGAAACCATATATCCTTGAAATTACGACTGCATCGTATACA
ATTATGATATAAATTAAAACATTAT

3' UTR

TTCTGGTCTCCACATATTACATATCAGCACCAATGGTTGAAAGGACATTGGCTTCT
TCTCTGGCAATGCATTCAATACAACATTGAAAATGACTTCAGCATATCAGTGTGCTTCGA
ACGTGTTCCGGAAGTACTCAAATGTGCTATGACTGAATTATTGTACATACATAACTTATTG
ATGTTCAATAATAATTGAAACGAAAAAAAAAAAAAA

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Figure 7

Derived primary structure of HtH2

Domain b

HRLFVTQVEDALIRRGSPIGVPYWDWTQPMALPGLADNATYRDPISGDSRHNPFHDVEVA
FENGRTERHPDSRLFEQPLFGKHTRLFDSIVYAFEQEDFCDFEVQFEMTHNNIHAWIGGGE
KYSMSSLHYTAFDPIFYLRHSNTDRLWAIWQALQIRRNRPYKAHCASSEERQPLKPFAFSS
PLNNNEKTYENSVPTNVYDYEGVLGYTYDDLNFGGMDLGQLEYYIQRQRQRDRTFAGFFLS
HIGTSANVEIIIDHGTLHTSVGTFAVLGGEKEMKWGFDRLYKYEITDELRLQNLRAADDVFS
ISVKVTDVDGSELSELIPSAIIIFERSH

Domain c

IDHQDPHDTIIRKNVDNLTPPEINSLRRAMADLQSDFKTAGGFQQIAAFHGEPKWCPSD
EKKFSCCVHGMVFPHWHRLTVQGENALRKHGCLGALPYWDWTRPLSHLPDLVLVSSRTT
PMPYSTVEARNPWYSGHIDTVGVDTTRSVRQELYEAPGFGHYGVAKQVLLALEQDDFCDF
EVQFEIAHNFIAHALVGGSEPYGMAISLRYTTYDPIFYLHHSNTDRLWAIWQALQYRGKPYN
SANCAIASMRKPLQPFGLTDEINPDDETROHAPFVFDYKNNFNYEYDTLDFNGLSISQL
DRELSRRKSHDRVFAGFLLHGIQQSALVKFFVCKSDDCDHYAGEFYILGDEAEMPWGYDR
LYKYEITEQLNALDLHIGDRFFIRYEAFDLHGTSLGSNIFPKPSVIHDEGA

Domain d

GHHQADEVVTAASHIRKNLKDLISKGEVESLRSAFLQLQNDGVYENIAKFHGKPGCLDD
NGRKVACCVHGMPTFPQWHLRYVLQVENALLERGSAVSVPYWDWTETFTELPSLIAEATYF
NSRQQTFDPNPFFRGKISFENAVTRDPQPELYVNRRYYQNVMVLFEQDNYCDFEIQFEMV
HNVLHAWLGG GRATYSISSLDYSAFDPVFFLHANTDRLWAIWQELQYRKKPYNEADCAIN
LMRKPLHPFDNSDLNHDPVTFKYSKPTDGFQYQNNFGYKYDNLEFNHFSIPRLEIIRI
RQDRVFAGFLLHNIGTSATVEIFVCVPTTSGEQNCENKAGTFAVLGGETEMAFHFDRLYRF
DISETLRDLGIQLDSHDFDLSIKIQGVNGSYLDPHILPEPSLIFVPGSS

Domain e

SFLRPDGHSDDILVRKEVNSLTRETASLIHALKSMQEDHSPDGFQAIASFHALPPLCSP
SAAHRYACCVHGMATFPQWHLRYTVQFQDALRRHGATGVPYWDWLRPQSHLPELVTMETY
HDIWSNRDFPNPFYQANIEFEGENITTEREVIADKLFVKGGHVFDKLVLQTSHP
SAEQENYCDFEIQFEILHNGVHTWVGGSRRTYSIGHLYAFYDPLFYLHHQTDRI
WAIWQELQEQRGLSGDEAHCALEQMREPLKPFSGAPYNWNQLTQDFSRPEDT
FDYRKGYFYEDNLEFLGMSVAELDQYIIIEHQENDRVFAGFLLSGFGGSAS
VNQVCRADSTCQDAGYFTVLGGS AEMA WAFD RLYKYDITETLEKML
RYDFTISVSLTANNGTVLSSLIPTPSVI FQRGH

Domain f

RDINTRSMSPNRVRRELSDLSARDLSSLKSA
LRLQEDDGPN
GYQALAAFHGLPAGCHDSR
GNEIACCIHGMPTFPQWHLRYTLQLE
MALRRHGSSVAI
PYWDWTKPISE
LPSLFTSPEYD
PWHD
AVVNNPFSKG
FVKFANTYTV
RDPQEMLFQLCE
HGESILYEQ
TLLALEQ
TDYCD
FEVQ
FEV
LHN
VIH
YLV
GGR
QTY
YAL
SSL
HYA
SYD
PFFF
FIH
HSF
V
DKM
W
V
W
QAL
OKR
RKL
PYK
RAD
CAV
NL
M
TKP
MRP
FDS
DMN
QNP
FTK
MHA
VPNT
LYD
YLY
SYD
NLE
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GRN
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IDR

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SRSHDRVAGFLLRGIGTSADVRFWICRNENDCHRGGIIFILGGAKEMPWSFDRNFKFDIT
HVLENAGISPEDVDAEFPYIKVEIHAVNKTMIPIPSSVIPAPTIIYSPGE

Domain g

GRAADSAHSANIAGSGVRKDVTTLTVSETENLRQALQGVIDDTGPNGYQAIASFHGSPPMC
EMNGRKVACCAHGMASFPHWHRLYVKQMEDALADHGSFIGIPYWDWTTAFTELPALVTDSE
NNPFHEGRIDHLGVTSRS PRDMLFNDPEQSESFFYRQVLLALEQTDYCQFEVQFELTHN
AIHSWTGGRSPYGMSTLEFTAYDPLFWLHHSNTDRIAVWQALQKYRGLPYNEAHCEIQVL
KQPLRPFNDDINHNPITKTNARPIDSFDYERFNYQYDTLSFHGKSIPELNDLLEERKREER
TFAAFLLRGIGCSADVVFDICRPNGDCVFAGTFAVLGGELEMPWSFDRLFRYDITRVMNQL
HLQYDSDFSFRVKLVATNGTELSSDLLKSPTIEHEL

Domain h

GAHRGPVEETEVTRQHTDGNAHFHRKEVDSLDEANNLKNALYKLQNDHSLTGYEAISGY
HGYPNLCEEGLDKIPLLPRMGIFPYWHRLTIQLERALEHNGALLGVPYWDWNKDLSSL
PAFFSDSSNNNPYFKYHIAGVGHDTVREPTSLIYNQPQIHGYDYLYYLALTTLEENNYWDF
EVQYEILHNAVHSWLGGSQYSMSTLEYSAFDPVFMILHSGLDRLWIIWQELQKIRRKPYN
FAKCAYHMMEEPLAPFSYPSINQDEFTRANSKPSTVFDSHKFGYHYDNLNRGHSIQELNT
IINDLRNTDRIYAGFVLSGIGTSASVKIYLRTDDNDEEVGTFVLGGEREMPWAYERVFKY
DITEVADRLKIKLWGHPLTSGTGDHILTNGIGGKQEPQTQILSSSTDLPIMTTMFLLSQXGR
NLHIPPKVVKKGTRIEFHPVDDSVTRPVVDLGSYTALFNCVVPFTYHGFELNHVYSVKP
GDYYVTGPTRDLCQNADVRHIHVEDE

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Figure 8

KLH1 cDNA sequence and intron structure

Domain b

GGCCTACCGTACTGGACTGGACTGAACCCATGACACACATTCGGGCTGGCAGGAAACA
AAACTTATGTGGATTCTCATGGTGATCCACACAAATCCTTTCATAGTCAGTGATTGC
ATTGAAAGAAAATGCTCCCCACACCAAAAGACAAATAGATCAAAGACTCTTAAACCCGCT
ACCTTGACACCACACAGACCTGTTCAACCAGATTGTATGCCCTTGAACAAGAAGATT
ACTGTGACTTGAAGTCCAATTGAGATTACCCATAACACGATTACGCTTGGACAGGAGG
AAGCGAACATTCTCAATGCGTCCCTACATTACACAGCTTCGATCCTTGTACTT
CACCAATTCTAACGTTGATCGTCTTGGGCCCTTGGCAAGCCTTACAGATGAGACGGCATA
AACCTACAGGGCCCACTGCGCCTATCTGGAACATATGCATCTGAAACCATTGCCCT
TTCATCTCCCCTTAACAATAACGAAAAGACTCATGCCATGCCATGCCAAACAAGATCTAC
GAATGAAAATGTCCTCATTACACATACGAAGATTAAACATTGGAGGCATCTCTGG
AAAACATAGAAAAGATGATCCACGAAAACCAGCAAGAAGACAGAAATATGCCGGTTTCT
CCTGGCTGGCATACTGACTTCAGCAAATGTTGATATCTTCATTAAAACCTACCGATTCCGTG
CAACATAAGGCTGGAACATTGCACTGCTCGGTGGAAGCAAGGAAATGAAGTGGGATTG
ATCGCGTTTCAAGTTGACATCACGCACGTTTGAAAGATCTCGATCTCACTGCTGATGG
CGATTTCGAAGTTACTGTTGACATCACTGAAGTCGATGGAACAAACTTGCATCCAGTCTT
ATTCCACATGCTCTGTCAATTGAGCATGCACGTGGTAAGCTGAATAGAG

Intron b/c

TTTTGTAATAATTATGAGAATTCTTACCTCAGAATAAGATGAGGTACATGGGTTTG
CAAAACTATTACGTTGAAATTAAATTAAATAACCGGACCCCTCCACTGGTACATATTAT
CTTTATAACGATAATAGCGATGATGATGATGATGATGATGATGATGATGATGATGATG
ATGATGCCGGTATTGACGTAATCCAGCGAATTAGATGACACCCTAAGGGTGCAGAAAGT
ATAaCAATTAGATTGCGTTGCATCTGTGTATGCGTGTGCTTAaCCAAAGTCAAATAA
AAAGTGCACACCCTAGTTATTGATAGAGCCTTACGATAAGAACAAATGTAATAA
ATTAGAACATAACTGAAACCTCCGAAAGAAGGGCTTTGTCAAGAGAGGTATCGACATGA
TTGACTTATAAACCTGTGCTCTATATTGGAACTGTCCACTTCTTGTGTGACTG
TAATCACATCGCACTATGGCTGCAAGACGTGTACGAGTACACTATATACTTACCTAATGAC
CAACCACAAGGCTGGCTTGTAAATTGTTATTGACAGAAATAAACACAGAATTCCAGC
ATTGGCTGGTGTATTAGCAAAACACCGATATGACACTCATGTTATTACATTTC
AG

Domain c

TTAAATTGACAAAGTGCCAAGGAGTCGTTATTGAAAAAAATGAGACCGTTGAGCCC
CGAGGAGATGAATGAACTTCGTAAGCCCTAGCCTACTGAAAGAGGACAAAGTGCCGGT
GGATTCAGCAGCTGGTGCATTCCATGGGAGCCAAATGGTGTCTAGTCCCGAAGCAT
CTAAAAAAATTGCTGCTGTGTTCA CGGCATGTCTGTGTTCCCTCACTGGCATCGACTGTT
GACGGTTAGAGTGAAGGAAATGCTTGTGAGACGACATGGCTACGATGGAGCTTGGCGTACTGG
GATTGGACCTCTCTTAAACCTCCGAACTGGCAGATCATGAGAAGTACGTCGACC
CTGAAGATGGGTAGAGAACATAACCTGGTGTGATGGTATAGATACAGTCGACAA
AACAAACAAGAACAGTGTGAGAACAGCCTGAGTTGGTATTGATGACATTACAG
AGCATTGCCAAACAAGTACTGCTAGCGTTGAAACAGGACAAATTCTGTGACTTGAATTC
AATATGAGATTGCCATAACTACATCCATGCACTTGTAGGAGGGCGTCAGCCTTATGGTAT
GGCATCGCTTCGCTACACTGCTTTGATCCACTATTCTACTTGCATCACTCTAAACAGAT
CGTATATGGCAATATGGCAGGCTTACAGAAGTACAGAGGAAACCGTACAACGTTGCTA
ACTGTGCTGTTACATCGATGAGAGAACCTTGCAACCATTGGCCTCTGCCAATATCAA
CACAGACCATGTAACCAAGGAGCATTGCCCCATTCAACGTTTTGATTACAAGACCAAT

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TTCAATTATGAATATGACACTTGAATTAAACGGTCTCTCAATCTCTCAGTTGAATAAAA
AGCTCGAAGCGATAAAGAGCCAAGACAGGTCTTGCAGGCTCCTGTTATCTGGTTCAA
GAAATCATCTCTGTTAAATTCAATATTGCACCGATAGCAGCAACTGTCACCCGCTGGA
GAGTTTACCTCTGGGTGATGAAAACGAGATGCCATGGGCATACTGATAGAGTCTTCAAAT
ATGACATAACCGAAAAACTCCACGATCTAAAGCTGCATGCAGAAGACCACTTCTACATTGA
CTATGAAGTATTGACCTAAACAGCAAGCCTGGAAAAGATTGTTCAAGCAGCCTTC
GTCATTGAAACCAAGAATAG

Intron c/d

GTACTTGTATATGTTCGAATATTGCCGATACCTTCAATATATACCTTATCAAAGTAA
TTGATTAATCTGAAGTAATTTCTTCCAGTAGAGATTAGCTGATACAACAAGAATTG
CCCTGTTGTATGTCACTTATTTCATCAAACGATTGAGTGGAGCTGTCCATGCCACAAT
GGGGTCTCTGTAACCTCTCGTATGGGTATAGATTATAGACGTGGCAGACCTACGTA
TAACTAATATTTGTGTAATGTCGTTCAAG

Domain d

GTCACCAGTGAAGGCAGACTATCAAGCTGAAGTAACCTCTGCCAACCGTATCGAAAAAA
CATTGAAAATCTGAGCCTGGTGAACCTGAAAGTCTGAGAGCTGCCTTCCTGGAAATTGAA
AACGATGAACTTACGAATCAATAGCTAAATTCCATGGTAGCCCTGGTTGTGCCAGTTAA
ATGGTAACCCCATCTCTTGTGTCCATGCCATGCCACTTCCCTCACTGGCACAGACT
GTACGTGGTTGTCGTTGAGAATGCCCTCTGAAAAAAGGATCATCTGTAGCTGTTCCCTAT
TGGGACTGGACAAAACGAATCGAACATTACCTCACCTGATTTCAGACGCCACTTACTACA
ATTCCAGGCAACATCACTATGAGACAAACCCATTCCATCATGGCAAAATCACACACGAGAA
TGAAATCACTACTAGGGATCCCAAGGACAGCCTCTCCATTCACTACTACTTTACGAGCAG
GTCCTTACGCCTGGAGCAGGATAACTTCTGTGATTGAGATTGAGATATTAC
ACAATGCATTGCAATTCTTACTTGGTGGCAAAGGTAATATTCCATGTCAAACCTTGATTA
CGCTGCTTTGATCCCTGTGTTCTCCTCATCACGCCAACGACTGACAGAAATCTGGCAATC
TGGCAAGACCTTCAGAGGTTCCGAAAACGCCATACCGAGAAGCGAATTGCGCTATCCAAT
TGATGCACACGCCACTCCAGCCGTTGATAAGAGCGACAACAATGAGCAGGGCAACGAAAAC
GCATGCCACTCCACATGATGGTTGAATATCAAACAGCTTGGTTATGCTTACGATAAT
CTGGAACCTGAATCACTACTCGATTCCCTCAGCTGATCACATGCTGCAAGAAAGAAAAAGGC
ATGACAGAGTATTGCTGGCTTCCTCACAATATTGGAACatCTGCCGATGCCATGT
ATTTGTATGTCTCCAACTGGGAACACACGAAGGACTGCAGTCAGTGGCTGTATGTC
TCCATCTAGGCAGTCAAACGGAGATGTCCTTGATTTGACAGACTTACAAACTTGACA
TAACTAAAGCCTGAAAAAGAACGGTGTGCACCTGCAAGGGATTGATCTGGAAATTGA
GATTACGGCTGTGAATGGATCTCATCTAGACAGTCATGTCATCCACTCTCCACTATACTG
TTTGAGGCCGGAACAG

Intron d/e

GTAACATTGTCAGTAAACCAACACTGCAGTCTATTGCAATTACGATAATAACAA
TTTTGAAATATCTTATTAAAGCAAAGGTTCTAGAGACAAACAGCCGGCTCTAATTAA
TTTTTGCACCTACGCTTGAGTAAAGATCTGCAAATGGCAACCCCTACCTATACTATTAA
AATATAATGTTACATTGTTAATAACACTCATATTGTCAG

Domain e

ATTCTGCCACACAGATGATGGACACACTGAACCAAGTGAATTGCAACCTGGCT
ATTGGACAAAGCGTCAACAACTGTCAGTGGTGAAGGCCCTCGAGTCCATGAAAGCCGACCAT
TCATCTGATGGGTCCAGGCAATCGCTTCCATGCTCTCCTCTTGTCCATCAC
CAGCTGCTCAAAGAGGTTGCGTGCGTCCATGGCATGCCAACCTCCGCAATG

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Figure 9

Derived primary structure of KLH1

Domain b

GLPYWDWTEPMTHIPGLAGNKTIVDHSHGASHTNPFHSSVIAFEENAPHTKRQIDQRLFKPA
TFGHHTDLFNQILYAFEQEDYCDFEVQFEITHNTIHAWTGGSEHFSMSSLHYTAFDPLFYF
HHSNVDRLWAVWQALQMRRHKPYRAHCAISLEHMHLKPFASFSSPLNNNEKTHANAMPNKIY
DYENVLHYTYEIDLTFGGISLENIEKMIHENQQEDRIYAGFLLAGIRTSANVDIFIKTTDSV
QHKAGTFAVLGGSKEMKGFDKFDITHVLKDLDLTADGDFEVTVVDITEVDGTLASSL
IPHASVIREHARGKLN

Domain c

VKFDKVPRSRLIRKNVDRILSPEEMNELRKALALLKEDKSAGGFQQLGAFHGEPKWCPSP
SKKFACCVHGMSPVPHWRLLTVQSENALRRHGYDGALPYWDWTSPLNHLPELADHEKYVD
PEDGVEKHNPFWDGHIDTVDKTTTRSVQNKLFEQPEFGHYSIAKQVLLALEQDNFCDFEI
QYEIAHNYIHALVGGAQPYGMASLRYTAFDPLFYLHHSNTDRIWAIWQALQKYRGKPYNVA
NCAVTSMREPLQPFGLSANINTDHVTKEHSPVFNVDYKTNFNYEYDTLEFNGLSISQLNK
KLEAIKSQDRFFAGFLLSGFKSSLVKFNICTDSSNCHPAGEFYLLGDENEMPWAYDRVFK
YDITEKLHDLKLHAEDHFYIDYEVFDLKPAISLGKDLFKQPSVIHEPRI

Domain d

GHHEGEVYQAEVTSANRIRKNIENSLGELESRAAFLEIENDGTYESIAKFHGSPGLCQL
NGNPISCCVHGMPTFPWHRLYVVVVENALLKKGSSVAVPYWDWTKRIEHLPHLISDATYY
NSRQHHYETNPFHGKITHENEITTRDPKDSLHFSDYFYEQVLYALEQDNFCDFEIQLEIL
HNALHSLLGGKGKYSMSNLDYAAFDPVFLHATTDRIWAIWQDLQRFRKRPYREANCAIQ
LMHTPLQPFDKSDNNDEATKTHATPHDGFEYQNSFGYAYDNLELNHYSIPQLDHMLQERKR
HDRVFAGFLLHNIGTSADGHVFVCLPTGEHTKDCSHEAGMFSILGGQTEMSFVFDRLYKLD
ITKALKKNGVHLQGDFDLEIEITAVNGSHLD SHVIHSPTILFEAG

Domain e

DSAHTDDGHTEPVMIRKDITQLDKRQQLSLVKALESMKADHSSDGFQAIASFHALPPLCPS
PAASKRFACCVHGMPTFPQWHRLYTVQFQDSLRKGAVVGLPYWDWTLP

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Figure 10

KLH2 cDNA sequence and intron structure

Domain b

GGCCTGCCCTACTGGGATTGGACCATGCCAATGAGTCATTTGCCAGAACTGGCTACAAGTG
AGACCTACCTCGATCCAGTTACTGGGAAACTAAAAACAAACCCCTTCATCACGCCCAAGT
GGCGTTGAAATGGTGTAAACAAGCAGGAATCCTGATGCCAACTTTTATGAAACCAACT
TACGGAGACCACACTTACCTCTTCGACAGCATGATCTACGCATTGAGCAGGAAGACTCT
GCGACTTGAAGTCCAATATGAGCTCACGCATAATGCAATACATGCATGGGTTGGAGGCAG
TGAAAAGTATTCAATGTCTCTTCACTacacTGCTTTGATCCTATATTACCTCCAT
CACTCAAATGTTGATCGTCTCTGGCCATTGGCAAGCTTCAAATCAGGAGAGGCAAGT
CTTACAAGGCCACTGCGCCTCGTCAAGAAAGAGAACATTAAAGCCTTGCATTCAAG
TTCCCCACTGAACAAACAACGAGAAAACGTACCACAACCTGTCCCCACTAACGTTATGAC
TATGTGGGAGTTTGCACTATCGATATGATGACCTTCAGTTGGCGGTATGACCATGTCAG
AACTTGAGGAATATATTACAAGCAGACACAACATGATAGAACCTTGCAGGATTCTCCT
TTCATATATTGGAACATCAGCAAGCGTAGATATCTTCATCAATCGAGAAGGTATGATAAA
TACAAAGTGGGAAGTTTGTAGTACTTGGGGATCCAAAGAAATGAAATGGGGCTTGATA
GAATGTACAAGTATGAGATCACTGAGGCTCTGAAGACGCTGAATGTTGCACTGGATGATGG
GTTCACTGAGGATCTGAGATCACCGATGTTGATGGATCTCCCCATCTGAGATCTCATT
CCACCTCCTGCTATAATCTTGaACGTGGTCaTG

Intron 2b/c

AGGTATTTAAAAAGTAATAAAACCaTATTTGAATGCGCTTATGAAATATCGTGTGAC
TGGTTCTTAGTTACATGGAGTGTAAACAACATGCTCCATCAGTTGACATATACTGCTCAC
ACAAAGTAAGGGATATTGATAATGATAACAAATATAATCAAAGCGGTATACTATCAAGA
CTTATTACACATAATTACAGGTGAAGGGAGGTGTATCGTCTACTGATCAGGTGAGGCC
AGAGAAGTCCCAGTTGAGTCTTGCAGAAGATGATGTTAGGCATGGGTCGAATCACCAA
ATCACATGACTTCAATAACGGTTGGACCACCTCGAGCAGATGCAAGCAGTAGAGCGT
CTACGCATGCTCCTGATAAGGCACCAATCTGTTCTGGGAATCAGtCGCCACTCCTCT
GTAGTGCCACGCTCATTTCTGCTACGGTCTGGTACCTGCTATCGGgtCTTGATCCGTAT
CCCAAGGATGTCCACACATGTTCAAGGTGAGAGGTGGGGACATCGCTGGCCACGGTaa
GGtCTGAATTGATGCCGTGAAAGTGAGCTCTGACAAACtGAGCATGGtGAGCTCTGACG
TTGTCGTCTGAAAGATGAATCAGCTcCaTGAACAGCGAGCAAAGGGCAGGACGTGTTGGT
CAATGCAGTTGTCCTGCAGTACACACCTGTCACTCGCCACTACAAGCGTGTAGATCTGT
ACGACCAGTCATGGAGATCCCAGCCACATCATAACGGACCCCTATCCATACCGATCATGA
GCCACCATAGCAGCGTCTGATGACGTTCTCCCTGTCGCTCGACATCCTcACACGGCCAA
AAGGAACGTGGACTCGTCACTGAACATGACATTAGCCAACCTGGCACTGTCACCGCTGA
TGTGGCGAGACCATTCCAGTCGAGCTTCTGGTGTCTGGCTTCATCGATAACACGACGT
AAGGTCTCGGGCGTGCAGACGGCTCTATGCAGGGATTTCGGATTGTCTGGGTGCTAAC
TCTGATCCCAGGTGCCTGCTGAAGTTGATGCTGGATCTGTGTGGCATTGAGATGGCGATT
CTTAGGACTGTGGAGATGATGAATCGATCTGACTTATGGTGGTGCACATTAGGACGTGCGG
TTCGTGTCTATCTGCACCTTCCAGTTGTCGGTGCAGCCTGTCGGTACCCGGCTGATTAC
TGACTGAGAATATCCATCTGCCGTGCGACATGAGCCTGTTGGCCACGCTGAAGCATTG
CAATGCCAGAGACGCTCTCAAAAGTCATTGACGATGGtTTCTGTTCAACAAATGACA
GCGTAAACAGtTTTGGtGCTTTATGCTTCCAAAGAGCATGAAAAACACGTTCTATgGG
TCGtGCACACCTTACATGACAAGtGtGAAAGtGACTTGcACCCCTTGTGtGtGTTGGATG
CACACTCTGTTACGTACTGATGCGATTGGCGTCTAAACATGTTTGGCGTCTAAACATG
TTTCCTGCTGATGATTCAATACTATTGTCATATTCTGGCATCAAACCAAACGAGT
AAATATATTCAATATCCCTACTTGTGTGAGTAGTATGATCACTGCAGACAACATATA

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GACAAtGCAgtTaCaCCGTCAACAATCCCAGTCATTAATTATGATGaCaCTTCCACACATA
GIGTCAGTGATTGTAATTCAaCTGTACACACTTTCCCGTAAACATTAGGATCTATATGA
CTAAATATATAACATTAGTATACGTGCAGTTGTATCGCTACGACATTGTTGTAACCTT
TGTTAATCATTaACAG

Domain c

CTGATGCCAAAGaCTTTGgCCATAGCAGAAAAATCAGgAAAGcCGTTGATTcTcTGACAGT
CgAAGAACAAAcTTCGTTGAGgCGAGcTATGgCAGATcTACAGGACGACAAACATCAGGG
GGTTCCAGCAGATTGCAGCATTCCACGGAGAACCAAAATGGTGTCCAAGCCCCgAAGCGG
AGAAAAAAATTGCATGCTGTTCATGGAATGGCTGTTTCCCTCACTGGCACAGATTGCT
GACAGTTCAAGGAGAAAATGCTCTGAGGAAACATGGCTTACTGGTGGACTGCCCTACTGG
GACTGGACTCGATCAATGAGGCCCTCCACATTTGTTGCTGATCCTACTTACAATGATG
CTATTCAGCCAGGAAGAAGATAACCCATGGCATCATGGTCACATAGACTCTGTTGGGCA
TGATACTACAAGAGAGATGTGCGTGATGATCTTATCAATCTCCTGGTTTCGGTCACTACACA
GATATTGCACAACAAGTCCTCTGGCCTTGAGCAGGACAGTTCTGTGATTTGAGGTAC
AATTGAAATTGCCATAATTACATGCACTGATTGGTGGTAACGAACCATAAGTAT
GTCATCTTGAGGTATACTACATACGATCCAATCTCTTGCACCACTCCAGTACAGAC
CGACTTTGGGCCATCTGGCAAGCAATCACTAGTGCAGGCCCTGCAGGTCGACCATAAGGG
AGAGCTCCCAACGCgtTGGAtGCAATCT

Domain g

ATGGCTGTGTTCCGCACTGGCACAGACTGTTGTGAAACAGATGGAGGACGCACTTGCTG
CTCATGGAGCTCATATTGGCATACCATCTGGGATTGGACAAGTGCCTTAGTCATCTGCC
CGCCCTAGTGACTGACCACGAGAACAAATCCCTCCACAC

Intron g (2)

GTATGTGTCAAATCGTTAGGAACACTGCCTTATCCATTAACTACGAGTACAAATGA
AAACGGAAACTGTGTGACCTCGAAAAGTGCATCTAAAGGATGCAATGTACACAATAAA
ATGCTCCGATAAAAGCGATGGCTAGAAATCATTTCCCTCTAAATTCCCTTCACACAGCT
CGGTTCGTTTAAGTAGGAACAAGTCTCTGCAAAAACATCAAAATAAGAGAACACAGAA
AAAACCTCATTCTGTTCTGTATTCCGAAAATGAAATTACAATTCTTCATTATAG

Domain g

GGCCATATTGGTCATCTGAATGTGGATACATCTGATCTCCAAGAGACATGCTGTTAATG
ATCCTGAACAAGGCTCAGAATCATTCTCTACAGACAGGTTCTTGACTCTAGAACAGAC
AGACTTCTGCCAATTGAAAGTTCAAGTTGACTTACACACAATGCCATCCACTCTGGACT
GGAGGACATACTCCATATGGAATGTCATCACTGGAATATACAGCATATGATCCACTCTTT
ATCTCCACCATTCCAACACTGATCGTATCTGGGCCATCTGGCAGGCACTCCAGAAAATATAG
AGGTCTTCCATACAACGCACTGCACTGCGATATCcaagttctgaaacaacctcTTAACCA
TTCAGCGAGTCCAGGAATCAAACCCAGTCACCAGAGCCAATTCTAGGGCCGTTGATTCAT
TTGATTATGAGAAATTCAATTATGACACACTTACCTCCACGGACTTTCTATCCC
AGAACTTGTATGCCATGCTCAAGAGAGAAAGAAGGAAGAGAGAACATTGCAAGCCTTCCTG
TTGCACGGATTGGCGCCAGTGCTGATGTTGTTGATGTCACACCTGATGGTCATT
GTGCCTTGCTGGAACCTCGCGGTACTTGGTGGGGAGCTTGAGATGCCCTGGCTTGA
AAGATTGTTCCGTTACGATATCACAAAGGTTCTCAAGCAGATGAATCTCACTATGATTCT
GAGTTCCACTTGAGTTGAAGATTGTTGGCACAGATGGAACAGAACTGCCATGGATCGTA
TCAAGAGCCCTACCATTGAACACCATGGAGGAG

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Intron g/h

GTATTTGAGATCCACATAATCTTCTACCCCTGTCTCATTTCTAATGCTCTTCAATACAC
AATTTATATAGCCTTGAGCTTCAGATGTATTACGGACAGGCATTACAGTATACATGTAAT
ATGGTTTCTGCTATTGCAAAATTGTGTCCTATCTGTTAGATCATGGCGGTGAC
CACCTAG

Domain h

GTCACGATCACAGTGAACGTCACGATGGATTTCAGGAAGGAAGTCGGTCCCTGTCCCT
GGATGAAGCCAATGACCTTAAAATGCACTGTACAAGCTGCAGAATGATCAGGGTCCAAT
GGATATGAATCAATAGCCGGTTACCATGGCTATCCATTCTGCCCTGAACATGGTGAAG
ACCAGTACGCATGCTGTGTCACGGAATGCCGTATTCCACATTGGCACAGACTTCATAC
AATCCAGTTGAGAGAGCTCTCAAAGAACATGGTTCTATTGGGTCTGCCATACTGGGAC
TGGAC

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Figure 11

Derived primary structure of KLH2

Domain b

GLPYWDWTMPMSHILPELATSETYLDPTGETKNNPFHHAQVAFENGVTSRNPDAKLFMKPT
YGDHTYLFDMSIYAFEQEDFCDFEVQYELTHNAIHAWVGGSEKYSMSSLHYTAFDPIFYLH
HSNVDRWLWAIWQALQIRRGKSYKAHCASSQEREPLKPFASFSSPLNNNEKTYHNSVPTNVYD
YVGVLHYRYDDLQFGGMTMSELEYYIHKQTQHDRTFAGFFLSYIGTSASVDIFINREGHDK
YKVGSFVVLGGSKEMKWGFDRMYKYEITEALKTLNAVDDGFSITVEITDVGSPPSADLI
PPPAIIFERGHA

Domain c

DAKDFGHSRKIRKAVDSLTVEEQTSLRRAMADLQDDKTSGGFQQIAAFHGEPKWCPSPPEAE
KKFACCVHGMAVFPWHRLLTQGENALRKHGFTGGLPYWDWTRSMSALPHFVADPTYNDA
ISSQEEDNPWHGHIDSVGHDTRDVRDDLYQSPGFGHYTDIAQQVLLAFEQDSFCDFEVQ
FEIAHNFIHALIGGNEPYSMSLRYTTYDPIFFLHSSTDRLWAIWQALQKYRGKPYNTAN
CAIASMRKPLQPFGLDSVINPDDETREHSVPFRVFDYKNNFDYEYESLAFNGLSIAQLDRE
LQRRKSHDRVFAGFLLHEIGQSAKHNVSDCDHYAGEFYILGDEAEMPWRYDRVYKYEITQQ
LHDDLHVGDNFFLKYEAFLNNGSLGGSIFSQPSVIPEPAAGMF

Domain d

GSHQADEYREAVTSASHIRKNIRDSEGEIESIRSAFLQIQKEGIYENIAKFHGKPGLEH
DGHPVACCVHGMPTFPWHRLYVLQVENALLERGSAVAVPYWDWTLPR

Domain g

MAVFPHWRLFVKQMEDALAAGAHIGIPYWDWTSAFSHLPALVTDHENNPFHGHIGHLN
VDTSRSPRDMLFNDPEQGSESFFYRQVLLTLEQTDQFCQFEVQFELTHNAIHSWTGGHTPYG
MSSLEYTAYDPLFYLHHSNTDRIWAIWQALQKYRGLPYNAAHCDIQVLKQPLKPFSESRN
NPVTRANSRAVDSFDYEKFNYQYDTLTFHGLSIPELDAMLQERKKEERTFAAFLLHGFGAS
ADVSFDVCTPDGHCAFAGTFAVLGGELEMPWSFERLFRYDITKVLKQMNLYDSEFHFE
IVGTDGTELPSDRIKSPTIEHHGG

Domain h

GHDHSERHDGFFRKEVGSLSLDEANDLKNALYKLQNDQGPNGYESIAGYHGYFLCPEHGE
DQYACCVHGMMPVFPWHRLHTIQFERALKEHGSHLGLPYWDW